

Molecular genetics

G. Ushakiran ^{1,*}, S. Ganga Sai Pradeepa ², S. Lavanya ², T. Sahithi Priya ² and P Krishna Kumari ²

¹ Department of pharmacology NRI college pharmacy Pothavarappadu Agiripalli, Andhra Pradesh, India.

² B. Pharmacy students NRI college Pharmacy Pothavarappadu, Andhra Pradesh, India.

World Journal of Advanced Research and Reviews, 2023, 20(03), 1035–1039

Publication history: Received on 31 October 2023; revised on 13 December 2023; accepted on 15 December 2023

Article DOI: <https://doi.org/10.30574/wjarr.2023.20.3.2057>

Abstract

Molecular genetics, is the study of the biochemical mechanisms of It is the study of the biochemical nature of the genetic material and it's control of phenotype. It is the study of the connection between genotype and phenotype the connection was a chemical one. Molecular genetics often applies an "investigative approach" to determine the structure and function of genes in an organism's genome using genetic screens. Molecular genetics is a powerful methodology for linking mutations to genetic conditions that may aid the search for treatments for various genetics diseases. This field has provided insights into various biological processes, including gene regulation, heredity, evolution, and disease development. Molecular genetics techniques, such as PCR, DNA sequencing, and gene editing, have revolutionized our ability to manipulate and study genes. The integration of molecular genetics with other disciplines, like genomics and proteomics, has paved the way for advancements in personalized medicine and biotechnology .It involves analyzing the DNA and RNA molecules to understand how genetic information is stored, replicated, and expressed.

Keywords: Molecular Genetics; Phenotype; Genotype; Genetics; Biotechnology; Replicated; Mutations; Integration; Revolutionized Polymerase Chain Reaction

1. Introduction

Molecular genetics or molecular biology is the study of bio chemical mechanisms heritage. It's the study of the biochemical nature of the inheritable material and its control of phenotype.it is the study of the connection between genotype and phenotype (1) The connection is a chemical one. control of phenotype is the one the two places of DNA (recap). You have formerly been exposed to the conception of the central dogma of Molecularbiology.is that the connection between genotype and phenotype is DNA (genotype) to RNA to enzyme to cell chemistry to phenotype. James Watson and Francis crick entered the 1953 Nobel prize for the discovery of the structure of the DNA patch (2). This is the alternate most important discovery in the history of the biology. Ranking just behind Charles Darwin. This discovery marked in the morning of an violent study of molecular biology bone that dominates ultramodern biology that will continue to do so into foreseeable future The essential characteristics of molecular Genetics is that gene products are studied through the genes that render them(3) this contrasts with a biochemical approach in which the gene product has a part in the process that are studying in vivo, but it cure'nt inescapably tell how direct that part is bio chemistry by discrepancy, tells what a factor can do in vitro but it cure n't inescapably mean that it does in vivo. The inheritable and biochemical approaches tell you different effects

Genetics has a role but not how direct Biochemistry what protein can do in vitro but not whether it really does in vivo These approaches therefore different things both are needed and are equally valuable. when one can combine these approaches to the figure out what a xvi gene/ protein does the resulting conclusions are much stronger than if only one use one of the strategies (4)

* Corresponding author: G. Ushakiran.

2. Types of modern method of genetics

2.1. Genome wide association studies

Genome-wide association studies (GWAS) have become a powerful tool in uncovering the genetic factors contributing to complex traits, including facial features⁹(5). By analyzing the genomes of thousands or even millions of individuals, GWAS identifies common genetic

variants associated with specific facial characteristics. These studies have provided valuable insights into the genes and genetic pathways involved in shaping facial morphology (6)

2.2. Three -dimensional face imaging

Three-dimensional (3D) facial imaging has revolutionized the field of facial genetics by enabling accurate and detailed measurements of facial structures (7) Utilizing advanced imaging technologies, such as 3D scanners or photogrammetry, researchers can capture high-resolution images of individuals' faces. These images are then analyzed using sophisticated software to extract precise measurements and create detailed facial models for genetic analysis (8)

2.3. Facial land marking and shape analysis

Facial landmarking involves identifying specific points on the face, such as the corners of the eyes or the tip of the nose, to map facial features accurately (9). Combining facial landmarking with shape analysis methods, researchers can quantify and compare facial shapes and identify subtle differences. These techniques help identify specific facial features that may be influenced by genetic variations (10)

2.4. Whole exome sequencing (WES) and whole genome sequencing (WGS)

Whole exome sequencing (WES) and whole genome sequencing (WGS) are advanced genetic technologies that provide comprehensive information about an individual's genetic makeup. WES focuses on sequencing the protein-coding regions of the genome, while WGS encompasses the entire genome (11). By analyzing the genetic data obtained through these techniques, researchers can identify rare genetic variants associated with facial features and gain insights into the underlying genetic architecture (12).

2.5. Functional genomics

. Functional Genomics: Functional genomics involves studying the function and activity of genes within the context of facial development. By investigating how specific genes are expressed during facial development, researchers can uncover their roles in shaping facial features. Techniques such as RNA sequencing and chromatin profiling provide valuable information about gene expression patterns and regulatory mechanisms involved in facial development. (13)

2.6. Integration of genetics and facial recognition technology:

Recent advancements in facial recognition technology have opened up exciting possibilities for studying the genetic basis of facial features. By combining genetic data with facial recognition algorithms, researchers can establish correlations between specific genetic variants and facial characteristics (14) This integration allows for large-scale studies involving thousands of individuals, facilitating the identification of genetic factors influencing facial morphology (15)

3. The modern methods of genetics of current aspects

Genetics, the study of heredity and the variation of inherited traits, has witnessed a revolution in recent decades with the advent of modern technologies and methodologies. This article delves into the current aspects of genetics, highlighting the breakthroughs and advancements that have transformed our understanding of the genetic code and its applications in various fields (16)

3.1. Genomic sequencing and precision medicine:

One of the most significant developments in genetics is the widespread adoption of genomic sequencing. The Human Genome Project, completed in 2003, marked a major milestone in deciphering the entire human genetic code. Since then, the cost of sequencing has dramatically reduced, enabling large-scale genomic studies and personalized medicine.

Researchers can now analyse an individual's genetic makeup to predict disease susceptibility, tailor treatments, and even develop targeted therapies for specific genetic conditions (17)

3.2. Crispr-cas9 and genome editing

The emergence of CRISPR-Cas9 gene editing technology has revolutionized the field of genetics. CRISPR-Cas9 allows scientists to precisely edit DNA sequences, enabling targeted modifications and corrections in the genetic code (18) This breakthrough technique has the potential to cure genetic disorders, enhance crop yields, and eradicate certain diseases by modifying the genes of disease-causing organisms. However, ethical concerns and careful regulation are crucial as the technology continues to advance (19)

3.3. Gene therapy and genetic medicine

Gene therapy, the process of introducing healthy genes into cells to treat genetic diseases, has shown tremendous promise in recent years (20). With the development of viral vectors and gene delivery systems, scientists can target and replace faulty genes, offering potential cures for previously incurable conditions. Gene therapy has demonstrated success in treating disorders such as spinal muscular atrophy and inherited retinal diseases, paving the way for a new era of genetic medicine (21).

3.4. Pharmacogenomics and drug development

Pharmacogenomics explores the relationship between an individual's genetic makeup and their response to drugs. By analysing genetic variations, researchers can predict an individual's drug metabolism, efficacy, and potential side effects (22) This knowledge allows for personalized drug prescriptions, optimizing treatment outcomes and reducing adverse reactions. Pharmaceutical companies are increasingly incorporating genomic data in drug development processes to identify suitable patient populations and design targeted therapies (23)

3.5. Agricultural genetics and crop improvement

Modern genetics has revolutionized agricultural practices, enhancing crop yield, quality, and resistance to diseases. Through genetic engineering and marker-assisted breeding, scientists can introduce desirable traits into plants, leading to improved productivity and sustainability (24) Genetic modification has allowed the development of crops with increased nutritional value, drought tolerance, and pest resistance, addressing global food security challenges (25)

3.6. Genetic data and privacy concerns

As genetic testing becomes more accessible and affordable, concerns regarding genetic data privacy have gained prominence. Genetic information is highly personal and sensitive, containing valuable insights into an individual's health and ancestry (26) Striking a balance between the benefits of genetic research and ensuring privacy and security is crucial for maintaining public trust and ethical standards (27)

4. The Modern Method of Genetics: Exploring Further Aspects

The field of genetics has undergone significant advancements in recent years, fueled by modern methodologies and technologies. This article explores additional aspects of genetics, highlighting the latest breakthroughs and their impact on various domains. From epigenetics and non-coding RNAs to synthetic biology and gene drives, these emerging areas are shaping the future of genetics research (28)

4.1. EPIGENETICS: Unveiling Gene Regulation Beyond the DNA Sequence

Epigenetics investigates heritable changes in gene expression patterns that do not involve alterations in the DNA sequence itself. It focuses on modifications to DNA and its associated proteins, known as histones, that influence gene activity. Epigenetic mechanisms, such as DNA methylation and histone modifications, play a crucial role in development, aging, and disease. Understanding these processes can provide insights into complex traits and facilitate the development of targeted therapies (29)

4.2. NON-CODING RNAs: Unveiling the Functional Complexity of the Genome

Non-coding RNAs (ncRNAs) are RNA molecules that do not code for proteins but perform crucial regulatory functions in the cell. They participate in gene silencing, chromatin re modelling, and post-transcriptional regulation (30) Examples of ncRNAs include microRNAs, long non-coding RNAs, and circular RNAs. Research in this area has revealed

their involvement in development, disease progression, and response to therapy, offering new therapeutic targets and diagnostic markers (31)

4.3. Synthetic biology: engineering novel genetic systems

Synthetic biology combines biology, engineering, and computer science to design and construct new biological parts, devices, and systems. Researchers can engineer genetic circuits, modify existing organisms, and create entirely synthetic organisms with desired functionalities (32) This field holds potential for various applications, including biofuel production, bioremediation, and drug synthesis, as well as advancements in medical diagnostics and treatment (33)

4.4. Gene drives: altering inherited traits in populations

Gene drives are genetic systems that can promote the inheritance of specific traits throughout a population. They involve the biased inheritance of genes to increase their spread rapidly. Gene drives have the potential to alter or eradicate disease vectors, control invasive species, and address agricultural challenges. However, the development and deployment of gene drives raise ethical and ecological considerations that require careful assessment (34)

5. Conclusion

The modern method of genetics has brought about remarkable advancements across various fields, from personalized medicine and gene therapy to agriculture and drug development. Genomic sequencing, CRISPR-Cas9 gene editing, and pharmacogenomics have transformed our understanding of genetics and opened up new avenues for targeted treatments. While these advancements hold immense potential, careful ethical considerations and privacy regulations are necessary to ensure responsible and equitable use of genetic information. As technology continues to progress, genetics will continue to shape the future of healthcare, agriculture, and our understanding of life itself.

The modern method of genetics continues to advance with emerging aspects such as epigenetics, non-coding RNAs, synthetic biology, and gene drives. These areas provide a deeper understanding of regulation, functional complexity of the genome, engineered biological systems, and population-level genetic modifications. They hold significant potential for diverse applications in medicine, agriculture, and environmental conservation. However, along with these advancements, careful consideration of ethical, social, and ecological implications is crucial to ensure responsible and beneficial use of these technologies. The future of genetics research and its applications is promising, offering unprecedented opportunities to address complex challenges and improve our lives.

Compliance with ethical standards

Disclosure of conflict of interest

No conflict of interest to be disclosed.

Reference

- [1] Bonner j.1965 The molecular Biology of department oxford university press Fogle T1990 :Are the genes units of inheritance Biology
- [2] Sarkar S2000 information in genetics and developmental Biology
- [3] Waters c.k 1994 genes made molecular philosophy of science.
- [4] LewisCM 2002 Genetic association studies design analysis and interpretation
- [5] Price AL patternsom NJ Plenge RM Weinblatt Me shadick NA Prinicipal componets anlysis corrects for Startification in genome wide association studies
- [6] Webber RL Horton RA Tyndall Da Ludlow JB Tuned Aperture computed Tomography Theory and application for three -dimensional imaging
- [7] Plooi JM Maal TJ Haers P Borstalap Wa Kuijpers -Jagatam AM Berges SJ .Digital three dimensional image fusion processes for planning and evaluating
- [8] Wu Y JI.Q.facial landmark detection a literearture survey 2019
- [9] Kemelmachar _Shlizerman I.Basri R 3D face reconstruction from a single imagre using a single reference shape

- [10] Wetterstrand K. DNA Sequencing costs data from the NHGRI Genome sequencing programme
- [11] Teer JK, Mullikin JC Exome sequencing of the sweet spot before whole genomes. *Hum Mol genetics*
- [12] Bentley, D. R., Balasubramanian, S., Swerdlow, H. P., Smith, G. P., Milton, J., Brown, C. G., Hall, K. P., Evers, D. J., Barners, C. L., Bignell, H., R. et al. (2008) accurate whole human genome sequencing using reversible terminator chemistry *nature*, 456, 53-59.
- [13] Lopez Ar Giro-i-Nieto X Burdick J Marques O. Skinlesion classification from dermoscopic images using deep learning techniques. 3
- [14] Loos HS Wieczkoerck D Wiirtz RP Von der malsburg C Horsthemke B Computer based Recognition Of dysmorphic faces
- [15] Venter, J. C., Adams, M. D., Myers, E. W., Li, P. W., Mural, R. J., Sutton, G. G., ... & Holt, R. A. (2001). The sequence of the human genome. *Science*, 291(5507), 1304-1351.
- [16] Collins, F. S., & Varmus, H. (2015). A new initiative on precision medicine. *New England Journal of Medicine*, 372(9), 793-795.
- [17] Doudna, J. A., & Charpentier, E. (2014). The new frontier of genome engineering with CRISPR-Cas9. *Science*, 346(6213)
- Mingozzi, F., Bennicelli, J., ... & Testa, F. (2008). Safety and efficacy of gene transfer for Leber's congenital amaurosis. *New England Journal of Medicine*, 358(21), 2240-2248.
- [18] Baltimore, D., Berg, P., Botchan, M., Carroll, D., Charo, R. A., Church, G., ... & Wu, F. (2015). A prudent path forward for genomic engineering and germline gene modification. *Science*, 348(6230), 36-38.
- [19] Maguire, A. M., Simonelli, F., Pierce, E. A., Pugh Jr, E. N., Mingozzi, F., Bennicelli, J., ... & Testa, F. (2008). Safety and efficacy of gene transfer for Leber's congenital amaurosis. *New England Journal of Medicine*, 358(21), 2240-2248.
21. Mendell, J. R., Al-Zaidy, S., Shell, R., Arnold, W. D., Rodino-Klapac, L. R., Prior, T. W., ... & Kaspar, B. K. (2017). Single-dose gene-replacement therapy for spinal muscular atrophy. *New England Journal of Medicine*, 377(18), 1713-1722.
- [20] Relling, M. V., & Evans, W. E. (2015). Pharmacogenomics in the clinic. *Nature*, 526(7573), 343-350.
- [21] Innocenti, F., & Ratain, M. J. (2011). Pharmacogenetics of anticancer agents: lessons from amonafide and irinotecan. *Drug Metabolism and Disposition*, 39(4), 631-635.
- [22] Paine, J. A., Shipton, C. A., Chaggar, S., Howells, R. M., Kennedy, M. J., Vernon, G., ... & Lawton, M. A. (2005). Improving the nutritional value of Golden Rice through increased pro-vitamin A content. *Nature Biotechnology*, 23(4), 482-487.
- [23] Jones, J. D., & Dangl, J. L. (2006). The plant immune system. *Nature*, 444(7117), 323-329
- [24] McGuire, A. L., Fisher, R., Cusenza, P., Hudson, K., Rothstein, M. A., & McGraw, D. (2008). Confidentiality, privacy, and security of genetic and genomic test information in electronic health records: points to consider. *Genetics in Medicine*, 10(7), 495-499.
- [25] Budin-Ljøsne, I., & Harris, J. R. (2010). A clash of principles: Genetic information and the need for an adequate ethical and legal framework. *Journal of Law, Medicine & Ethics*, 38(2), 336-345.
- [26] Bird, A. (2007). Perceptions of epigenetics. *Nature*, 447(7143), 396-398.
- [27] Feinberg, A. P., & Irizarry, R. A. (2010). Evolution in health and medicine Sackler colloquium: Stochastic epigenetic variation as a driving force of development, evolutionary adaptation, and disease. *Proceedings of the National Academy of Sciences*, 107(Supplement 1), 1757-1764.
- [28] Bartel, D. P. (2009). MicroRNAs: target recognition and regulatory functions. *Cell*, 136(2), 215-233.
- [29] Hobom, B. Surgery of genes at the doorstep of synthetic biology. *medizin* 14-21
- [30] Rawls R Synthetic biology makes its them debut chen. *Eng news* 49-53
- [31] Khalil, A. S., & Collins, J. J. (2010). Synthetic biology: applications come of age. *Nature Reviews Genetics*, 11(5), 367-379.
- nome regulation by long noncoding RNAs. *Annual Review of Biochemistry*, 81, 145-166.
- [32] Esvelt, K. M., Smidler, A. L., Catteruccia, F., & Church, G. M. (2014). Concerning RNA-guided gene drives for the alteration of wild populations. *eLife*, 3, e03401.
- Noble, C., Olejarz, J., Esvelt, K. M., & Church, G. M. (2019)